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<110> Barber, Brian Berinstein, Neil Moingeon, Philippe Tartaglia, James

TECH CENTER 1600/2900

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<120> Method of Inducing and/or Enhancing an Immune Response
      to Tumor Antigens
<130> 11014-15
<140> 09/693,754
<141> 2000-10-20
<150> 60/223,325
<151> 2000-08-07
<150> 60/160,879
<151> 1999-10-22
<160> 113
<170> PatentIn Ver. 2.0
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Ser Tyr Val Pro Leu Ala His Ser Ser Ser Ala Phe Thr Ile Thr
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His Phe Leu Arg Asn Gln Pro Leu Thr Phe Ala Leu Gln Leu His
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Gln Pro Leu Thr Phe Ala Leu Gln Leu His Asp Pro Ser Gly Tyr
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<211> 15
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Ser Thr Gly Leu Ile Ser Arg Ala Leu Val Val Thr His Thr Tyr
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Ser Arg Ala Leu Val Val Thr His Thr Tyr Leu Glu Pro Gly Pro
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Val Thr His Thr Tyr Leu Glu Pro Gly Pro Val Thr Ala Gln Val
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Ile Pro Leu Thr Ser Cys Gly Ser Ser Pro Val Pro Gly Thr Thr
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Val Pro Gly Thr Thr Asp Gly His Arg Pro Thr Ala Glu Ala Pro
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Asp Gly His Arg Pro Thr Ala Glu Ala Pro Asn Thr Thr Ala Gly
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Glu Val Val Gly Thr Thr Pro Gly Gln Ala Pro Thr Ala Glu Pro
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 Thr Pro Gly Gln Ala Pro Thr Ala Glu Pro Ser Gly Thr Thr Ser
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 Pro Thr Ala Glu Pro Ser Gly Thr Thr Ser Val Gln Val Pro Thr
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 <211> 15
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Val Gln Val Pro Thr Thr Glu Val Ile Ser Thr Ala Pro Val Gln
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<400> 61
Thr Glu Val Ile Ser Thr Ala Pro Val Gln Met Pro Thr Ala Glu
                  5
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<400> 62
Thr Ala Pro Val Gln Met Pro Thr Ala Glu Ser Thr Gly Met Thr
                   5
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 <400> 63
 Met Pro Thr Ala Glu Ser Thr Gly Met Thr Pro Glu Lys Val Pro
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 <211> 15
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Ser Thr Gly Met Thr Pro Glu Lys Val Pro Val Ser Glu Val Met
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Pro Glu Lys Val Pro Val Ser Glu Val Met Gly Thr Thr Leu Ala
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Val Ser Glu Val Met Gly Thr Thr Leu Ala Glu Met Ser Thr Pro
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Gly Thr Thr Leu Ala Glu Met Ser Thr Pro Glu Ala Thr Gly Met
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 <400> 68
 Glu Met Ser Thr Pro Glu Ala Thr Gly Met Thr Pro Ala Glu Val
 <210> 69
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Pro Glu Gly Pro Asp Ala Ser Ser Ile Met Ser Thr Glu Ser Ile
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Ala Ser Ser Ile Met Ser Thr Glu Ser Ile Thr Gly Ser Leu Gly
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Ser Thr Glu Ser Ile Thr Gly Ser Leu Gly Pro Leu Leu Asp Gly
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 Thr Gly Ser Leu Gly Pro Leu Leu Asp Gly Thr Ala Thr Leu Arg
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Leu Val Lys Arg Gln Val Pro Leu Asp Cys Val Leu Tyr Arg Tyr
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 Val Leu Tyr Arg Tyr Gly Ser Phe Ser Val Thr Leu Asp Ile Val
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Gln Gly Ile Glu Ser Ala Glu Ile Leu Gln Ala Val Pro Ser Gly
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Ala Glu Ile Leu Gln Ala Val Pro Ser Gly Glu Gly Asp Ala Phe
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 Ala Val Pro Ser Gly Glu Gly Asp Ala Phe Glu Leu Thr Val Ser
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Cys Gln Gly Gly Leu Pro Lys Glu Ala Cys Met Glu Ile Ser Ser
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Pro Lys Glu Ala Cys Met Glu Ile Ser Ser Pro Gly Cys Gln Pro
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Met Glu Ile Ser Ser Pro Gly Cys Gln Pro Pro Ala Gln Arg Leu
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Gln Leu Val Leu His Gln Ile Leu Lys Gly Gly Ser Gly Thr Tyr
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Ala Gly Leu Gly Gln Val Pro Leu Ile Val Gly Ile Leu Leu Val
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Tyr Arg Arg Leu Met Lys Gln Asp Phe Ser Val Pro Gln Leu
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 <210> 104
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Met Lys Gln Asp Phe Ser Val Pro Gln Leu Pro His Ser Ser Ser
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Pro His Ser Ser Ser His Trp Leu Arg Leu Pro Arg Ile Phe Cys
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Pro Arg Ile Phe Cys Ser Cys Pro Ile Gly Glu Asn Ser Pro Leu
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tat cca gag tg Tyr Pro Glu Tr 50	g aca gaa p Thr Glu	gcc cag a Ala Gln A 55	aga ctt Arg Leu	gac tgc tgg Asp Cys Trp 60	aga ggt Arg Gly	ggt 192 Gly							
caa gtg tcc ct Gln Val Ser Le 65	c aag gtc u Lys Val 70	agt aat o	gat ggg Asp Gly	cct aca ctg Pro Thr Leu 75	att ggt Ile Gly	gca 240 Ala 80							
aat gcc tcc tt Asn Ala Ser Ph	c tct att e Ser Ile 85	gcc ttg a	aac ttc Asn Phe 90	cct gga agc Pro Gly Ser	caa aag Gln Lys 95	gta 288 Val							
ttg cca gat gg Leu Pro Asp Gl	y Gln Val	Ile Trp	gtc aac Val Asn 105	aat acc atc Asn Thr Ile	atc aat Ile Asn 110	ggg 336 Gly							
agc cag gtg to Ser Gln Val Tr 115	g gga gga p Gly Gly	cag cca Gln Pro 120	gtg tat Val Tyr	ccc cag gaa Pro Gln Glu 125	. Thr Asp	gat 384 Asp							
gcc tgc atc tt Ala Cys Ile Ph 130	c cct gat e Pro Asp	ggt gga Gly Gly 135	cct tgc Pro Cys	cca tct ggc Pro Ser Gly 140	tct tgg Ser Trp	tct 432 Ser							
cag aag aga ag Gln Lys Arg Se 145	c ttt gtt r Phe Val 150	Tyr Val	tgg aag Trp Lys	acc tgg ggc Thr Trp Gly 155	caa tac Gln Tyr	tgg 480 Trp 160							
caa gtt cta g Gln Val Leu G	gg ggc cca y Gly Pro 165	gtg tct Val Ser	ggg ctg Gly Leu 170	agc att ggg Ser Ile Gly	g aca ggc Thr Gly 175	agg 528 Arg							
gca atg ctg g Ala Met Leu G 1	gc aca cac Ly Thr His 30	acg atg Thr Met	gaa gtg Glu Val 185	act gtc tac Thr Val Tyr	cat cgc His Arg 190	cgg 576 Arg							
gga tcc cgg a Gly Ser Arg S 195	gc tat gtg er Tyr Val	cct ctt Pro Leu 200	gct cat Ala His	tcc agc tca Ser Ser Ser 20!	r Ala Phe	acc 624 Thr							
att atg gac c Ile Met Asp G	ag gtg cct ln Val Pro	ttc tcc Phe Ser	gtg agc Val Ser	gtg tcc cas Val Ser Gl	g ttg cgg n Leu Arg	gcc 672 Ala							

210 215 220

	210					213					220					
ttg Leu 225	gat Asp	gga Gly	ggg Gly	aac Asn	aag Lys 230	cac His	ttc Phe	ctg Leu	aga Arg	aat Asn 235	cag Gln	cct Pro	ctg Leu	acc Thr	ttt Phe 240	720
gcc Ala	ctc Leu	cag Gln	ctc Leu	cat His 245	gac Asp	ccc Pro	agt Ser	ggc Gly	tat Tyr 250	ctg Leu	gct Ala	gaa Glu	gct Ala	gac Asp 255	ctc Leu	768
tcc Ser	tac Tyr	acc Thr	tgg Trp 260	gac Asp	ttt Phe	gga Gly	gac Asp	agt Ser 265	agt Ser	gga Gly	acc Thr	ctg Leu	atc Ile 270	tct Ser	cgg Arg	816
gca Ala	ctt Leu	gtg Val 275	gtc Val	act Thr	cat His	act Thr	tac Tyr 280	ctg Leu	gag Glu	cct Pro	ggc Gly	cca Pro 285	gtc Val	act Thr	gtt Val	864
cag Gln	gtg Val 290	gtc Val	ctg Leu	cag Gln	gct Ala	gcc Ala 295	att Ile	cct Pro	ctc Leu	acc Thr	tcc Ser 300	tgt Cys	ggc Gly	tcc Ser	tcc Ser	912
cca Pro 305	gtt Val	cca Pro	ggc Gly	acc Thr	aca Thr 310	gat Asp	gly ggg	cac His	agg Arg	cca Pro 315	act Thr	gca Ala	gag Glu	gcc Ala	cct Pro 320	960
aac Asn	acc Thr	aca Thr	gct Ala	ggc Gly 325	caa Gln	gtg Val	cct Pro	act Thr	aca Thr 330	gaa Glu	gtt Val	gtg Val	ggt Gly	act Thr 335	aca Thr	1008
cct Pro	ggt Gly	cag Gln	gcg Ala 340	cca Pro	act Thr	gca Ala	gag Glu	ccc Pro 345	tct Ser	gga Gly	acc Thr	aca Thr	tct Ser 350	gtg Val	cag Gln	1056
gtg Val	cca Pro	acc Thr 355	act Thr	gaa Glu	gtc Val	ata Ile	agc Ser 360	act Thr	gca Ala	cct Pro	gtg Val	cag Gln 365	atg Met	cca Pro	act Thr	1104
gca Ala	gag Glu 370	agc Ser	aca Thr	ggt Gly	atg Met	aca Thr 375	cct Pro	gag Glu	aag Lys	gtg Val	cca Pro 380	gtt Val	tca Ser	gag Glu	gtc Val	1152
atg Met 385	Gly	acc Thr	aca Thr	ctg Leu	gca Ala 390	gag Glu	atg Met	tca Ser	act Thr	cca Pro 395	gag Glu	gct Ala	aca Thr	ggt Gly	atg Met 400	1200
aca Thr	cct Pro	gca Ala	gag Glu	gta Val 405	tca Ser	att Ile	gtg Val	gtg Val	ctt Leu 410	tct Ser	gga Gly	acc Thr	aca Thr	gct Ala 415	gca Ala	1248
cag Gln	gta Val	aca Thr	act Thr 420	aca Thr	gag Glu	tgg Trp	gtg Val	gag Glu 425	acc Thr	aca Thr	gct Ala	aga Arg	gag Glu 430	cta Leu	cct Pro	1296
atc Ile	cct Pro	gag Glu 435	Pro	gaa Glu	ggt Gly	cca Pro	gat Asp 440	gcc Ala	agc Ser	tca Ser	atc Ile	atg Met 445	Ser	acg Thr	gaa Glu	1344
agt Ser	att Ile 450	aca Thr	ggt Gly	tcc Ser	ctg Leu	ggc Gly 455	ccc Pro	ctg Leu	ctg Leu	gat Asp	ggt Gly 460	Thr	gcc Ala	acc Thr	tta Leu	1392
agg Arg 465	Leu	gtg Val	aag Lys	aga Arg	caa Gln 470	Val	ccc Pro	ctg Leu	gat Asp	tgt Cys 475	Val	ctg Leu	tat Tyr	cga Arg	tat Tyr 480	1440

ggt tcc ttt tcc Gly Ser Phe Ser	gtc acc ctg Val Thr Leu 485	gac att gtc Asp Ile Val 490	cag ggt att ga Gln Gly Ile Gl	a agt gcc 1488 1 Ser Ala 495
gag atc ctg cag Glu Ile Leu Gln 500	gct gtg ccg Ala Val Pro	tcc ggt gag Ser Gly Glu 505	ggg gat gca tt Gly Asp Ala Ph 51	e Glu Leu
act gtg tcc tgc Thr Val Ser Cys 515	caa ggc ggg Gln Gly Gly	ctg ccc aag Leu Pro Lys 520	gaa gcc tgc at Glu Ala Cys Me 525	g gag atc 1584 t Glu Ile
tca tcg cca ggg Ser Ser Pro Gly 530	tgc cag ccc Cys Gln Pro 535	cct gcc cag Pro Ala Gln	cgg ctg tgc ca Arg Leu Cys Gl 540	g cct gtg 1632 n Pro Val
cta ccc agc cca Leu Pro Ser Pro 545	gcc tgc cag Ala Cys Gln 550	ctg gtt ctg Leu Val Leu	cac cag ata ct His Gln Ile Le 555	g aag ggt 1680 u Lys Gly 560
ggc tcg ggg aca Gly Ser Gly Thr	tac tgc ctc Tyr Cys Leu 565	aat gtg tct Asn Val Ser 570	ctg gct gat ac Leu Ala Asp Th	c aac agc 1728 r Asn Ser 575
ctg gca gtg gtc Leu Ala Val Val 580	agc acc cag Ser Thr Gln	ctt atc atg Leu Ile Met 585	cct ggt caa ga Pro Gly Gln Gl 59	u Ala Gly
ctt ggg cag gtt Leu Gly Gln Val 595	ccg ctg atc Pro Leu Ile	gtg ggc atc Val Gly Ile 600	ttg ctg gtg tt Leu Leu Val Le 605	g atg gct 1824 u Met Ala
gtg gtc ctt gca Val Val Leu Ala 610	tct ctg ata Ser Leu Ile 615	Tyr Arg Arg	aga ctt atg aa Arg Leu Met Ly 620	g caa gac 1872 s Gln Asp
ttc tcc gta ccc Phe Ser Val Pro 625	cag ttg cca Gln Leu Pro 630	cat agc agc His Ser Ser	agt cac tgg ct Ser His Trp Le 635	g cgt cta 1920 u Arg Leu 640
ccc cgc atc ttc Pro Arg Ile Phe	tgc tct tgt Cys Ser Cys 645	ccc att ggt Pro Ile Gly 650	gag aac agc co Glu Asn Ser Pr	c ctc ctc 1968 o Leu Leu 655
agt ggg cag cag Ser Gly Gln Gln 660	Val			1986
<210> 110 <211> 661 <212> PRT <213> Artificia	l Sequence			
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Ala Leu Leu Ala 20		Thr Lys Val 25	Pro Arg Asn G	n Asp Trp 0
Leu Gly Val Ser 35	Arg Gln Leu	a Arg Thr Lys 40	Ala Trp Asn As 45	g Gln Leu

Tyr Pro Glu Trp Thr Glu Ala Gln Arg Leu Asp Cys Trp Arg Gly Gly Gln Val Ser Leu Lys Val Ser Asn Asp Gly Pro Thr Leu Ile Gly Ala Asn Ala Ser Phe Ser Ile Ala Leu Asn Phe Pro Gly Ser Gln Lys Val Leu Pro Asp Gly Gln Val Ile Trp Val Asn Asn Thr Ile Ile Asn Gly Ser Gln Val Trp Gly Gly Gln Pro Val Tyr Pro Gln Glu Thr Asp Asp Ala Cys Ile Phe Pro Asp Gly Gly Pro Cys Pro Ser Gly Ser Trp Ser Gln Lys Arg Ser Phe Val Tyr Val Trp Lys Thr Trp Gly Gln Tyr Trp Gln Val Leu Gly Gly Pro Val Ser Gly Leu Ser Ile Gly Thr Gly Arg 170 Ala Met Leu Gly Thr His Thr Met Glu Val Thr Val Tyr His Arg Arg Gly Ser Arg Ser Tyr Val Pro Leu Ala His Ser Ser Ser Ala Phe Thr Ile Met Asp Gln Val Pro Phe Ser Val Ser Val Ser Gln Leu Arg Ala 215 Leu Asp Gly Gly Asn Lys His Phe Leu Arg Asn Gln Pro Leu Thr Phe Ala Leu Gln Leu His Asp Pro Ser Gly Tyr Leu Ala Glu Ala Asp Leu Ser Tyr Thr Trp Asp Phe Gly Asp Ser Ser Gly Thr Leu Ile Ser Arg Ala Leu Val Val Thr His Thr Tyr Leu Glu Pro Gly Pro Val Thr Val Gln Val Val Leu Gln Ala Ala Ile Pro Leu Thr Ser Cys Gly Ser Ser 295 Pro Val Pro Gly Thr Thr Asp Gly His Arg Pro Thr Ala Glu Ala Pro Asn Thr Thr Ala Gly Gln Val Pro Thr Thr Glu Val Val Gly Thr Thr 330 Pro Gly Gln Ala Pro Thr Ala Glu Pro Ser Gly Thr Thr Ser Val Gln Val Pro Thr Thr Glu Val Ile Ser Thr Ala Pro Val Gln Met Pro Thr 365 Ala Glu Ser Thr Gly Met Thr Pro Glu Lys Val Pro Val Ser Glu Val Met Gly Thr Thr Leu Ala Glu Met Ser Thr Pro Glu Ala Thr Gly Met 395

Thr Pro Ala Glu Val Ser Ile Val Val Leu Ser Gly Thr Thr Ala Ala 405 Gln Val Thr Thr Glu Trp Val Glu Thr Thr Ala Arg Glu Leu Pro 425 Ile Pro Glu Pro Glu Gly Pro Asp Ala Ser Ser Ile Met Ser Thr Glu Ser Ile Thr Gly Ser Leu Gly Pro Leu Leu Asp Gly Thr Ala Thr Leu Arg Leu Val Lys Arg Gln Val Pro Leu Asp Cys Val Leu Tyr Arg Tyr Gly Ser Phe Ser Val Thr Leu Asp Ile Val Gln Gly Ile Glu Ser Ala 485 Glu Ile Leu Gln Ala Val Pro Ser Gly Glu Gly Asp Ala Phe Glu Leu Thr Val Ser Cys Gln Gly Gly Leu Pro Lys Glu Ala Cys Met Glu Ile Ser Ser Pro Gly Cys Gln Pro Pro Ala Gln Arg Leu Cys Gln Pro Val 535 Leu Pro Ser Pro Ala Cys Gln Leu Val Leu His Gln Ile Leu Lys Gly Gly Ser Gly Thr Tyr Cys Leu Asn Val Ser Leu Ala Asp Thr Asn Ser Leu Ala Val Val Ser Thr Gln Leu Ile Met Pro Gly Gln Glu Ala Gly 585 Leu Gly Gln Val Pro Leu Ile Val Gly Ile Leu Leu Val Leu Met Ala Val Val Leu Ala Ser Leu Ile Tyr Arg Arg Arg Leu Met Lys Gln Asp 615 Phe Ser Val Pro Gln Leu Pro His Ser Ser Ser His Trp Leu Arg Leu 630 Pro Arg Ile Phe Cys Ser Cys Pro Ile Gly Glu Asn Ser Pro Leu Leu 650

Ser Gly Gln Gln Val 660

<210> 111

<211> 2106

<212> DNA

<213> Artificial Sequence

-<220>

<221> CDS

<222> (1)..(2106)

<2205

<223> Description of Artificial Sequence: modified CEA

<400> 11 atg gag Met Glu 1	tct c	cc t Pro S	tcg Ser 5	gcc Ala	cct Pro	ccc Pro	cac His	aga Arg 10	tgg Trp	tgc Cys	atc Ile	ccc Pro	tgg Trp 15	cag Gln	48
agg ctc Arg Leu	Leu L	etc a Leu :	aca Thr	gcc Ala	tca Ser	ctt Leu	cta Leu 25	acc Thr	ttc Phe	tgg Trp	aac Asn	ccg Pro 30	ccc Pro	acc Thr	96
act gcc Thr Ala	aag c Lys L 35	etc a Leu '	act Thr	att Ile	gaa Glu	tcc Ser 40	acg Thr	ccg Pro	ttc Phe	aat Asn	gtc Val 45	gca Ala	gag Glu	gly ggg	144
aag gag Lys Glu 50	gtg c Val L	ctt (Leu 1	cta Leu	ctt Leu	gtc Val 55	cac His	aat Asn	ctg Leu	ccc Pro	cag Gln 60	cat His	ctt Leu	ttt Phe	ggc Gly	192
tac agc Tyr Ser 65	tgg t Trp T	ac (Tyr :	aaa Lys	ggt Gly 70	gaa Glu	aga Arg	gtg Val	gat Asp	ggc Gly 75	aac Asn	cgt Arg	caa Gln	att Ile	ata Ile 80	240
gga tat Gly Tyr	gta a Val I	ata (gga Gly 85	act Thr	caa Gln	caa Gln	gct Ala	acc Thr 90	cca Pro	Gly ggg	ccc Pro	gca Ala	tac Tyr 95	agt Ser	288
ggt cga Gly Arg	Glu I	ata Ile 100	ata Ile	tac Tyr	ccc Pro	aat Asn	gca Ala 105	tcc Ser	ctg Leu	ctg Leu	atc Ile	cag Gln 110	aac Asn	atc Ile	336
atc cag Ile Gln	aat g Asn A	gac Asp	aca Thr	gga Gly	ttc Phe	tac Tyr 120	acc Thr	cta Leu	cac His	gtc Val	ata Ile 125	aag Lys	tca Ser	gat Asp	384
ctt gtg Leu Val 130	aat g Asn G	gaa Glu	gaa Glu	gca Ala	act Thr 135	ggc Gly	cag Gln	ttc Phe	cgg Arg	gta Val 140	tac Tyr	ccg Pro	gag Glu	ctg Leu	432
ccc aag Pro Lys 145	ccc t Pro S	tcc Ser	atc Ile	tcc Ser 150	agc Ser	aac Asn	aac Asn	tcc Ser	aaa Lys 155	ccc Pro	gtg Val	gag Glu	gac Asp	aag Lys 160	480
gat gct Asp Ala	gtg g Val A	Ala	ttc Phe 165	acc Thr	tgt Cys	gaa Glu	cct Pro	gag Glu 170	act Thr	cag Gln	gac Asp	gca Ala	acc Thr 175	tac Tyr	528
ctg tgg Leu Trp	Trp V	gta Val 180	aac Asn	aat Asn	cag Gln	agc Ser	ctc Leu 185	ccg Pro	gtc Val	agt Ser	ccc Pro	agg Arg 190	ctg Leu	cag Gln	576
ctg tcc Leu Ser	aat o Asn 0 195	ggc Gly	aac Asn	agg Arg	acc Thr	ctc Leu 200	act Thr	cta Leu	ttc Phe	aat Asn	gtc Val 205	aca Thr	aga Arg	aat Asn	624
gac aca Asp Thr 210	gca a Ala s	agc Ser	tac Tyr	aaa Lys	tgt Cys 215	gaa Glu	acc Thr	cag Gln	aac Asn	cca Pro 220	gtg Val	agt Ser	gcc Ala	agg Arg	672
cgc agt Arg Ser 225	gat t Asp :	tca Ser	gtc Val	atc Ile 230	ctg Leu	aat Asn	gtc Val	ctc Leu	tat Tyr 235	Gly	ccg Pro	gat Asp	gcc Ala	ccc Pro 240	720
acc att Thr Ile	tcc (Ser)	cct Pro	cta Leu 245	aac Asn	aca Thr	tct Ser	tac Tyr	aga Arg 250	tca Ser	Gly	gaa Glu	aat Asn	ctg Leu 255	aac Asn	768

ctc Leu	tcc Ser	tgc Cys	cac His 260	gca Ala	gcc Ala	tct Ser	aac Asn	cca Pro 265	cct Pro	gca Ala	cag Gln	tac Tyr	tct Ser 270	tgg Trp	ttt Phe	816
gtc Val	aat Asn	ggg Gly 275	act Thr	ttc Phe	cag Gln	caa Gln	tcc Ser 280	acc Thr	caa Gln	gag Glu	ctc Leu	ttt Phe 285	atc Ile	ccc Pro	aac Asn	864
atc Ile	act Thr 290	gtg Val	aat Asn	aat Asn	agt Ser	gga Gly 295	tcc Ser	tat Tyr	acg Thr	tgc Cys	caa Gln 300	gcc Ala	cat His	aac Asn	tca Ser	912
gac Asp 305	act Thr	ggc Gly	ctc Leu	aat Asn	agg Arg 310	acc Thr	aca Thr	gtc Val	acg Thr	acg Thr 315	atc Ile	aca Thr	gtc Val	tat Tyr	gag Glu 320	960
cca Pro	ccc Pro	aaa Lys	ccc Pro	ttc Phe 325	atc Ile	acc Thr	agc Ser	aac Asn	aac Asn 330	tcc Ser	aac Asn	ccc Pro	gtg Val	gag Glu 335	gat Asp	1008
gag Glu	gat Asp	gct Ala	gta Val 340	gcc Ala	tta Leu	acc Thr	tgt Cys	gaa Glu 345	cct Pro	gag Glu	att Ile	cag Gln	aac Asn 350	aca Thr	acc Thr	1056
tac Tyr	ctg Leu	tgg Trp 355	tgg Trp	gta Val	aat Asn	aat Asn	cag Gln 360	agc Ser	ctc Leu	ccg Pro	gtc Val	agt Ser 365	ccc Pro	agg Arg	ctg Leu	1104
cag Gln	ctg Leu 370	tcc Ser	aat Asn	gac Asp	aac Asn	agg Arg 375	acc Thr	ctc Leu	act Thr	cta Leu	ctc Leu 380	agt Ser	gtc Val	aca Thr	agg Arg	1152
aat Asn 385	gat Asp	gta Val	gga Gly	ccc Pro	tat Tyr 390	gag Glu	tgt Cys	gga Gly	atc Ile	cag Gln 395	aac Asn	gaa Glu	tta Leu	agt Ser	gtt Val 400	1200
gac Asp	cac His	agc Ser	gac Asp	cca Pro 405	gtc Val	atc Ile	ctg Leu	aat Asn	gtc Val 410	Leu	tat Tyr	ggc	cca Pro	gac Asp 415	gac Asp	1248
ccc Pro	acc Thr	att Ile	tcc Ser 420	Pro	tca Ser	tac Tyr	acc Thr	tat Tyr 425	Tyr	cgt Arg	cca Pro	ggg Gly	gtg Val 430	Asn	ctc Leu	1296
agc Ser	ctc Leu	tcc Ser 435	. CAs	cat His	gca Ala	gcc Ala	tct Ser 440	Asn	cca Pro	. cct Pro	gca Ala	cag Gln 445	Tyr	tct Ser	tgg Trp	1344
ctg Leu	att Ile 450	Asp	ggg Gly	aac Asn	atc Ile	cag Gln 455	Gln	cac His	aca Thr	caa Gln	gag Glu 460	тей	ttt Phe	atc : Ile	tcc Ser	1392
aac Asr 465	Ile	act Thr	gag Glu	g aag 1 Lys	aac Asn 470	Sex	gga Gly	cto Lev	tat Tyr	acc Thr 475	Cys	cag Glr	gcc Ala	aat Asn	aac Asn 480	1440
tca Ser	gcc Ala	agt Sei	ggo Gly	cac His	Ser	agg Arg	g act g Thi	aca Thr	gto Val 490	r ràs	g aca	a ato	aca Thr	yto Val 495	tct Ser	1488
gcg Ala	gag Glu	g ctg 1 Lei	g cco i Pro 500) Lys	g ccc s Pro	tco Sei	ato Ile	tco Ser 505	s Sei	c aac c Asr	aac Ası	tco n Sei	c aaa c Lys 510	S PIC	gtg Val	1536
gag	g gad	c aag	g gat	t gct	gtg	gco	c tto	aco	tgt	c gaa	a cci	gaq	g gct	cag	g aac	1584

Glu Asp Ly	ys Asp 15	Ala	Val	Ala	Phe 520	Thr	Cys	Glu	Pro	Glu 525	Ala	Gln	Asn	
aca acc ta Thr Thr Ty 530	ac ctg yr Leu	tgg Trp	tgg Trp	gta Val 535	aat Asn	ggt Gly	cag Gln	agc Ser	ctc Leu 540	cca Pro	gtc Val	agt Ser	ccc Pro	1632
agg ctg ca Arg Leu G 545	ag ctg ln Leu	tcc Ser	aat Asn 550	ggc Gly	aac Asn	agg Arg	acc Thr	ctc Leu 555	act Thr	cta Leu	ttc Phe	aat Asn	gtc Val 560	1680
aca aga aa Thr Arg A	at gac sn Asp	gca Ala 565	aga Arg	gcc Ala	tat Tyr	gta Val	tgt Cys 570	gga Gly	atc Ile	cag Gln	aac Asn	tca Ser 575	gtg Val	1728
agt gca aa Ser Ala A	ac cgc sn Arg 580	agt Ser	gac Asp	cca Pro	gtc Val	acc Thr 585	ctg Leu	gat Asp	gtc Val	ctc Leu	tat Tyr 590	Gly	ccg Pro	1776
gac acc co Asp Thr P	cc atc ro Ile 95	att Ile	tcc Ser	ccc Pro	cca Pro 600	gac Asp	tcg Ser	tct Ser	tac Tyr	ctt Leu 605	tcg Ser	gga Gly	gcg Ala	1824
gac ctc a Asp Leu A 610	ac ctc sn Leu	tcc Ser	tgc Cys	cac His 615	tcg Ser	gcc Ala	tct Ser	aac Asn	cca Pro 620	tcc Ser	ccg Pro	cag Gln	tat Tyr	1872
tct tgg c Ser Trp A 625	gt atc rg Ile	aat Asn	630 Gly ggg	ata Ile	ccg Pro	cag Gln	caa Gln	cac His 635	aca Thr	caa Gln	gtt Val	ctc Leu	ttt Phe 640	1920
atc gcc a Ile Ala L	aa atc ys Ile	acg Thr 645	cca Pro	aat Asn	aat Asn	aac Asn	ggg Gly 650	acc Thr	tat Tyr	gcc Ala	tgt Cys	ttt Phe 655	gtc Val	1968
tct aac t Ser Asn L	tg gct eu Ala 660	act Thr	ggc Gly	cgc Arg	aat Asn	aat Asn 665	tcc Ser	ata Ile	gtc Val	aag Lys	agc Ser 670	atc Ile	aca Thr	2016
gtc tct g Val Ser A 6	ca tct la Ser 75	gga Gly	act Thr	tct Ser	cct Pro 680	ggt Gly	ctc Leu	tca Ser	gct Ala	Gly Ggg	gcc Ala	act Thr	gtc Val	2064
ggc atc a Gly Ile M 690	tg att Met Ile	gga Gly	gtg Val	ctg Leu 695	gtt Val	Gly	gtt Val	gct Ala	ctg Leu 700	ata Ile	tag			2106
<210> 112 <211> 701 <212> PRT <213> Art	;	l Sed	quen	ce										
<400> 112 Met Glu S 1	er Pro	Ser 5	Ala	Pro	Pro	His	Arg 10	Trp	Cys	Ile	Pro	Trp 15	Gln	
Arg Leu L	eu Leu 20		Ala	Ser	Leu	Leu 25	Thr	Phe	Trp	Asn	Pro 30		Thr	
Thr Ala I	ys Leu 35	Thr	Ile	Glu	Ser 40	Thr	Pro	Phe	Asn	Val 45		Glu	Gly	,
Lys Glu V 50	/al Leu	Leu	Leu	Val 55		Asn	Leu	Pro	Gln 60		Leu	Phe	Gly	

Tyr Ser Trp Tyr Lys Gly Glu Arg Val Asp Gly Asn Arg Gln Ile Ile Gly Tyr Val Ile Gly Thr Gln Gln Ala Thr Pro Gly Pro Ala Tyr Ser Gly Arg Glu Ile Ile Tyr Pro Asn Ala Ser Leu Leu Ile Gln Asn Ile Ile Gln Asn Asp Thr Gly Phe Tyr Thr Leu His Val Ile Lys Ser Asp Leu Val Asn Glu Glu Ala Thr Gly Gln Phe Arg Val Tyr Pro Glu Leu 135 Pro Lys Pro Ser Ile Ser Ser Asn Asn Ser Lys Pro Val Glu Asp Lys Asp Ala Val Ala Phe Thr Cys Glu Pro Glu Thr Gln Asp Ala Thr Tyr Leu Trp Trp Val Asn Asn Gln Ser Leu Pro Val Ser Pro Arg Leu Gln 185 Leu Ser Asn Gly Asn Arg Thr Leu Thr Leu Phe Asn Val Thr Arg Asn Asp Thr Ala Ser Tyr Lys Cys Glu Thr Gln Asn Pro Val Ser Ala Arg Arg Ser Asp Ser Val Ile Leu Asn Val Leu Tyr Gly Pro Asp Ala Pro Thr Ile Ser Pro Leu Asn Thr Ser Tyr Arg Ser Gly Glu Asn Leu Asn Leu Ser Cys His Ala Ala Ser Asn Pro Pro Ala Gln Tyr Ser Trp Phe 265 Val Asn Gly Thr Phe Gln Gln Ser Thr Gln Glu Leu Phe Ile Pro Asn Ile Thr Val Asn Asn Ser Gly Ser Tyr Thr Cys Gln Ala His Asn Ser 295 Asp Thr Gly Leu Asn Arg Thr Thr Val Thr Thr Ile Thr Val Tyr Glu Pro Pro Lys Pro Phe Ile Thr Ser Asn Asn Ser Asn Pro Val Glu Asp 330 Glu Asp Ala Val Ala Leu Thr Cys Glu Pro Glu Ile Gln Asn Thr Thr 345 Tyr Leu Trp Trp Val Asn Asn Gln Ser Leu Pro Val Ser Pro Arg Leu Gln Leu Ser Asn Asp Asn Arg Thr Leu Thr Leu Leu Ser Val Thr Arg Asn Asp Val Gly Pro Tyr Glu Cys Gly Ile Gln Asn Glu Leu Ser Val Asp His Ser Asp Pro Val Ile Leu Asn Val Leu Tyr Gly Pro Asp Asp

415 410 405

Pro Thr Ile Ser Pro Ser Tyr Thr Tyr Tyr Arg Pro Gly Val Asn Leu 425 Ser Leu Ser Cys His Ala Ala Ser Asn Pro Pro Ala Gln Tyr Ser Trp Leu Ile Asp Gly Asn Ile Gln Gln His Thr Gln Glu Leu Phe Ile Ser 455 Asn Ile Thr Glu Lys Asn Ser Gly Leu Tyr Thr Cys Gln Ala Asn Asn Ser Ala Ser Gly His Ser Arg Thr Thr Val Lys Thr Ile Thr Val Ser Ala Glu Leu Pro Lys Pro Ser Ile Ser Ser Asn Asn Ser Lys Pro Val Glu Asp Lys Asp Ala Val Ala Phe Thr Cys Glu Pro Glu Ala Gln Asn Thr Thr Tyr Leu Trp Trp Val Asn Gly Gln Ser Leu Pro Val Ser Pro Arg Leu Gln Leu Ser Asn Gly Asn Arg Thr Leu Thr Leu Phe Asn Val Thr Arg Asn Asp Ala Arg Ala Tyr Val Cys Gly Ile Gln Asn Ser Val Ser Ala Asn Arg Ser Asp Pro Val Thr Leu Asp Val Leu Tyr Gly Pro 585 Asp Thr Pro Ile Ile Ser Pro Pro Asp Ser Ser Tyr Leu Ser Gly Ala Asp Leu Asn Leu Ser Cys His Ser Ala Ser Asn Pro Ser Pro Gln Tyr Ser Trp Arg Ile Asn Gly Ile Pro Gln Gln His Thr Gln Val Leu Phe Ile Ala Lys Ile Thr Pro Asn Asn Asn Gly Thr Tyr Ala Cys Phe Val Ser Asn Leu Ala Thr Gly Arg Asn Asn Ser Ile Val Lys Ser Ile Thr Val Ser Ala Ser Gly Thr Ser Pro Gly Leu Ser Ala Gly Ala Thr Val Gly Ile Met Ile Gly Val Leu Val Gly Val Ala Leu Ile

<210> 113

<211> 9

<212> PRT

<213> Artificial Sequence

<223> Description of Artificial Sequence: modified CEA epitope

<400> 113 Tyr Leu Ser Gly Ala Asp Leu Asn Leu 1 5